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Keiko Sato

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Toshihisa Ohtsuka

<120> mammalian Prickle gene

<130> E1-A0207P

<140> PCT/JP03/15223

<141> 2003-11-28

<150> JP 2002-346447

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<170> PatentIn Ver. 2.1

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Val His Gln Tyr Tyr Ser Cys Leu Pro Glu Glu Lys Val Pro Tyr Val
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Asn Ser Pro Gly Glu Lys Leu Arg Ile Lys Gln Leu Leu His Gln Leu
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Pro Pro His Asp Asn Glu Val Arg Tyr Cys Asn Ser Leu Asp Glu Glu
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Glu Lys Arg Glu Leu Lys Leu Phe Ser Asn Gln Arg Lys Arg Glu Asn
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Leu Gly Arg Gly Asn Val Arg Pro Phe Pro Val Thr Met Thr Gly Ala
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Glu Ala Glu Gly Arg His Trp His Met Arg His Phe Cys Cys Phe Glu 210 215 220

Cys Glu Thr Val Leu Gly Gly Gln Arg Tyr Ile Met Lys Glu Gly Arg
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Thr Pro Ser Leu Asn Arg Asp Pro Ile Trp Arg Ser Arg Asp Glu Pro
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Phe His Tyr Gly Asn Lys Met Glu Gln Asn Gln Ser Gln Ser Pro Leu

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Glu Gly Gly Ile Ser Thr Gln Gln Cys Arg Pro Arg Arg Pro Leu Ser
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Gly Ser Met Glu Ser Leu Ala Leu Ser Asn Ala Thr Gly Leu Ser Ala 545 550 555 560

Glu Gly Gly Ala Lys Arg Gln Glu His Leu Ser Arg Phe Ser Met Pro
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Met Gly Thr Leu Asn Ser Ser Met Gln Phe Arg Ser Ala Glu Ser Val
595 600 605

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Arg Met His Gln Ser Phe Asp Phe Asp Gly Gly Ile Ala Ser Ser Lys
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Leu Pro Gly Gln Glu Gly Val His Ile Gln Pro Met Ser Glu Arg Thr
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725 730 735

Glu Ser Met Gly Gln Gly Ser Arg Arg Asp Leu Tyr Ser Gln Cys Pro
740 745 750

Arg Thr Val Ser Asp Leu Ala Leu Gln Asn Ala Phe Gly Glu Arg Trp
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Gly Pro Tyr Phe Thr Glu Tyr Asp Trp Cys Ser Thr Cys Ser Ser Ser 770 780

Ser Glu Ser Asp Asn Glu Gly Tyr Phe Leu Gly Glu Pro Ile Pro Gln
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Pro Ala Arg Leu Arg Tyr Val Thr Ser Asp Glu Leu Leu His Lys Tyr

805 810 815

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Gln	Arg	Asn	Ser	Thr	Ser	Asp	Asp	Asp	Ser	Gly	Cys	Ala	Leu	Glu	Glu	
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tat	age	tac	ctc	cca	ซลล	മുമ	222	σ††	ccc	tat	gtc	aac	age	cct	gga	606
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70 75 80

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tgt	gga	ggt	cag	att	aag	ggc	ggc	gac	atc	gct	gtg	ttt	gca	tca	cgt	846
Cys	Gly	Gly	Gln	Ile	Lys	Gly	Gly	Asp	Ile	Ala	Val	Phe	Ala	Ser	Arg	
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Cys	Asn	Glu	Leu	Leu	Val	Asp	Leu	Ile	Tyr	Phe	Tyr	Gln	Asp	Gly	Lys	
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+					+-+	44.	.	44						4.4		1070
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His	Ala	Thr	Glu	Asn	Cys	Phe	Cys	Cys	Ala	His	Cys	Lys	Lys	Ser	Leu	
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Ala	Cys	Ser	Ala	Gly	Glu	Asp	Pro	Asn	Gly	Ser	Asp	Ser	Ser	Asp	Ser	
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Gly	Lys	Asn	Lys	Gly	Lys	Thr	Glu	Glu	Thr	Met	Leu	Asn	Gln	His	Ser	
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Gln	Leu	Gln	Val	Ser	Ser	Asn	Arg	Leu	Ser	Ala	Asp	Val	Asp	Pro	Leu	
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Ser	Val	Gln	Met	Asp	Leu	Leu	Ser	Leu	Ser	Ser	Gln	Thr	Pro	Ser	Leu	
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Asn Arg Asp Pro Ile Trp Arg Ser Arg Asp Glu Pro Phe His Tyr Gly

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Ala	Gln	Pro	Asp	Met	Trp	Ala	Lys	His	Phe	Ser	Asn	Pro	Lys	Arg	Ser	
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	gag															1854
Arg	Glu	Asp	Tyr	Tyr	Pro		Arg	Leu	Met	Ser		Glu	Ser	Tyr	Ser	
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	atg															1902
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aac	tcc	tca	atg	caa	ttc	agg	agc	gca	gag	tct	gtc	cgc	agc	cta	ctg	2238
Asn	Ser	Ser	Met	Gln	Phe	Arg	Ser	Ala	Glu	Ser	Val	Arg	Ser	Leu	Leu	

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Glu	Val	Ile	Ala	Arg	Leu	Lys	Asp	Arg	Pro	Pro	Leu	Arg	Ala	Arg	Glu	
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Gln	Gly	Ser	Arg	Arg	Asp	Leu	Tyr	Ser	Gln	Cys	Pro	Arg	Thr	Val	Ser	
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cga	tat	gtc	aca	agt	gat	gag	ctc	ctg	cac	aag	tac	agc	tct	tac	ggt	2862
Arg	Tyr	Val	Thr	Ser	Asp	Glu	Leu	Leu	His	Lys	Tyr	Ser	Ser	Tyr	Gly	

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Val Pro Lys Ser Ser Thr Leu Gly Gly Arg Gly Gln Leu His Ser Arg

825

830

835

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840 845

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35 40 45

Leu Gly Ser Gly Gly Met Gly Leu Pro Ser Ser Ser Ser Ala Ser Ala
50 55 60

Leu Tyr Ala Ala Gln Ala Ala Gly Ile Leu Pro Thr Ser Pro Leu
65 70 75 80

Pro Leu Gln Arg His Gln Gln Tyr Leu Pro Pro His His Gln Gln His

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Pro Gly Ala Gly Met Gly Pro Gly Pro Gly Ser Gly Ala Ala Ala Gly

100 105 110

Pro Pro Leu Gly Pro Gln Tyr Ser Pro Gly Cys Ser Ala Asn Pro Lys
115 120 125

Tyr Ser Asn Ala Gln Leu Pro Pro Pro Pro His His His Gln Leu
130 135 140

Ser Pro Ala Leu Ser Thr Pro Ser Pro Pro Ser Leu Leu His His Pro 145 150 155 160

Ala Gly Gly Thr Ser Ser Ala Ser Ala His Ala Pro Phe Leu Gly Gly

165 170 175

Pro His Met Asp Met Gln Arg Gln Ser His Ser Asp Asp Ser Gly
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Cys Ala Leu Glu Glu Tyr Thr Trp Val Pro Pro Gly Leu Arg Pro Asp 195 200 205

Gln Val Arg Leu Tyr Phe Ser Gln Ile Pro Asp Asp Lys Val Pro Tyr
210 215 220

Val Asn Ser Pro Gly Glu Gln Tyr Arg Val Arg Gln Leu Leu His Gln 225 230 235 240

Leu Pro Pro His Asp Asn Glu Val Arg Tyr Cys His Ser Leu Thr Asp

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Glu Glu Arg Lys Glu Leu Arg Leu Phe Ser Thr Gln Arg Lys Arg Asp
260 265 270

Ala Leu Gly Arg Gly Asn Val Arg Gln Leu Met Ser Ala Arg Pro Cys
275 280 285

Asp Gly Cys Asp Asp Leu Ile Ser Thr Gly Asp Ile Ala Val Phe Ala
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Thr Arg Leu Gly Pro Asn Ala Ser Trp His Pro Ala Cys Phe Ala Cys
305 310 315 320

Ser Val Cys Arg Glu Leu Leu Val Asp Leu Ile Tyr Phe His Arg Asp 325 330 335

Gly Arg Met Tyr Cys Gly Arg His His Ala Glu Thr Leu Lys Pro Arg

340 345 350

Cys Ser Ala Cys Asp Glu Ile Ile Leu Ala Asp Glu Cys Thr Glu Ala 355 360 365

Glu Gly Arg Ala Trp His Met Asn His Phe Gly Cys His Glu Cys Asp 370 375 380

Lys	Gln	Leu	Gly	Gly	Gln	Arg	Tyr	Ile	Met	Arg	Glu	Gly	Lys	Pro	Tyr
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Cys Leu His Cys Phe Asp Ala Met Phe Gly Glu Tyr Cys Asp Tyr Cys
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Gly Glu Gly Ile Gly Val Asp Gln Gly Gln Met Ser His Asp Gly Gln
420 425 430

His Trp His Ala Thr Asp Glu Cys Phe Ser Cys Asn Thr Cys Arg Cys
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Ser Leu Leu Gly Arg Ala Phe Leu Pro Arg Arg Gly Gly Ile Tyr Cys
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Ser Ile Ala Cys Ser Lys Gly Glu Pro Pro Thr Pro Ser Asp Ser Ser 465 470 475 480

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Pro His Pro Gln Ala Pro Leu Pro Ala Arg Ile Pro Ser Ser His Ala
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Ala 545					Asp 550	Gln	Ser	Lys	Ser	Tyr 555	Ala	Ala	Ser	Asp	Ser 560
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Gly	Asp	Leu	Thr 580	Asp	Phe	Ser	Gly.	Gly 585	Arg	Ala	Ser	Ser	Thr 590	Ser	G1r
Asn	Leu	Ser 595	Pro	Leu	Asn	Ser	Pro 600	G1y	Asp	Phe	Gln	Pro 605	His	Phe	Leu
Pro	Lys 610	Pro	Met	Glu	Leu	Gln 615	Arg	Asp	Gly	Val	Tyr 620	Asn	Phe	Asn	G1u
Met 625	Ser	Ser	Asn	Leu	Asp 630	Ala	Ala	Trp	Pro	Ala 635	Lys	Pro	Thr	Asn	Ser 640

Tyr His Leu Gln Arg Gln Leu Leu Glu Asn Pro His Thr Ala Ser Met

650

655

645

Pro Glu Leu Ala Gly Lys Leu Val Ala Pro Pro Ala His Met Gln His
660 665 670

Leu Ser Gln Leu His Ala Val Ser Ser His Gln Phe Gln Gln His Glu 675 680 685

Tyr Ala Asp Ile Val His Pro Pro Pro Pro Pro Gly Glu Ile Pro 690 695 700

Glu Leu Pro Thr Pro Asn Leu Ser Val Ala Ser Thr Ala Leu Pro Pro
705 710 715 720

Glu Leu Met Gly Ser Pro Thr His Ser Ala Gly Asp Arg Ser Leu Asn
725 730 735

Thr Pro Met Ser Thr Gln Ser Ala Ser His Ala Pro Pro His Pro Val
740 745 750

Ser Ile Leu Ser Gly Ala Ser Ser Ser Pro Met Ser Gly Glu Pro
755 760 765

Ala Lys Lys Gly Val Arg Phe Glu Gly Ile Pro Asp Thr Leu Pro
770 775 780

Arg Ser Arg Ser Tyr Ser Gly Asn Gly Ala Gly Thr Ser Ser Gly Gly
785 790 795 800

Glu Arg Glu Arg Asp Arg Asp Lys Asp Lys Glu Gly Gly Arg His
805 810 815

Gly His Gly His Ser Ser Arg Arg Arg Arg Arg Lys Ser Ser Ser 820 825 830

Ser Ser Ser His His Arg Ser Gly Ser Gly His Arg Ser His Ser Thr

835

840

845

Thr Arg Ala Asp Thr Tyr Ala Pro Ala Gln Pro Leu Ser Ser Tyr 850 855 860

Gln Gly Pro Pro Ser Val Leu Gln Ala Ala Asn Leu Val His Glu Ser 865 870 875 880

Pro Ser Arg Gln Gln Arg Glu Arg Glu Arg Glu Arg Glu Arg Lys Glu

885

890

895

Ser Glu Asp Tyr Met Met Met Tyr Gln Leu Pro Gln Arg Arg His Tyr
915 920 925

Gly Gly Val Arg Val Ser Tyr Val Pro Asn Asp Ala Leu Ala Tyr Asp

930 935 940

Arg Lys Arg Lys Pro Ser Glu Leu Gly Gly Asp Lys Asp Lys Asn Cys 945 950 955 960

Ile Ile Ser